

SEQUENCE LISTING

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Shirley, Bret A.
Bajwa, Kamaljit K.

<120> Stabilized FGF Formulations Containing
Reducing Agents

<130> PP16021.002

<150> 60/229,238

<151> 2000-08-31

<160> 8

<170> FastSEQ for Windows Version 4.0

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<211> 146

<212> PRT

<213> Homo sapiens

<400> 1

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Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp
          35           40           45
Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser
          50           55           60
Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly
65           70           75           80
Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu
          85           90           95
Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Thr
          100          105          110
Ser Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Ser
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Lys Thr Gly Pro Gly Gln Lys Ala Ile Leu Phe Leu Pro Met Ser Ala
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Lys Ser
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<210> 2

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<212> PRT

<213> Bos taurus

<400> 2

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Pro Ala Leu Pro Glu Asp Gly Gly Ser Gly Ala Phe Pro Pro Gly His
 1           5           10           15
Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu
          20           25           30

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Tyr Cys Lys Asn Gly Gly Phe Phe Leu Arg Ile His Pro Asp Gly Arg
 35 40 45
 Val Asp Gly Val Arg Glu Lys Ser Asp Pro His Ile Lys Leu Gln Leu
 50 55 60
 Gln Ala Glu Glu Arg Gly Val Val Ser Ile Lys Gly Val Cys Ala Asn
 65 70 75 80
 Arg Tyr Leu Ala Met Lys Glu Asp Gly Arg Leu Leu Ala Ser Lys Cys
 85 90 95
 Val Thr Asp Glu Cys Phe Phe Phe Glu Arg Leu Glu Ser Asn Asn Tyr
 100 105 110
 Asn Thr Tyr Arg Ser Arg Lys Tyr Ser Ser Trp Tyr Val Ala Leu Lys
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 Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu
 20 25 30
 cgc atc cac ccc gac ggc cga gtt gac ggg gtc cgg gag aag agc gac 144
 Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp
 35 40 45
 cct cac atc aag cta caa ctt caa gca gaa gag aga gga gtt gtg tct 192
 Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser
 50 55 60
 atc aaa gga gtg tgt gct aac cgt tac ctg gct atg aag gaa gat gga 240
 Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly
 65 70 75 80
 aga tta ctg gct tct aaa tgt gtt acg gat gag tgt ttc ttt ttt gaa 288
 Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu
 85 90 95
 cga ttg gaa tct aat aac tac aat act tac cgg tca agg aaa tac acc 336
 Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Thr
 100 105 110

agt tgg tat gtg gca ctg aaa cga act ggg cag tat aaa ctt gga tcc 384
 Ser Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Ser
 115 120 125

aaa aca gga cct ggg cag aaa gct ata ctt ttt ctt cca atg tct gct 432
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 Lys Ser *
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 Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu
 20 25 30

cga atc cac cca gat ggg cga gta gat ggg gta cga gaa aaa tcc gat 144
 Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp
 35 40 45

cca cac atc aaa cta caa cta caa gcc gaa gaa cga ggg gta gta tcc 192
 Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser
 50 55 60

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 Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly
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cga cta cta gcc tcc aaa tgt gta acc gat gaa tgt ttc ttc ttc gaa 288
 Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu
 85 90 95

cga cta gaa tcc aac aac tat aac acc tat cga tcc cga aaa tat tcc 336
 Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Ser
 100 105 110

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